

Leptin signaling regulates glucose homeostasis, but not adipostasis, in the zebrafish

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Leptin is the primary adipostatic factor in mammals. Produced largely by adipocytes in proportion to total adipose mass, the hormone informs the brain regarding total energy stored as triglycerides in fat cells. The hormone acts on multiple circuits in the brain to regulate food intake, autonomic outflow, and endocrine function to maintain energy balance. In addition to regulating adipose mass, mammalian leptin also plays a role in the regulation of glucose homeostasis and as a gating factor in reproductive competence. Leptin-deficient mice and people exhibit early onset profound hyperphagia and obesity, diabetes, and infertility. Although leptin and the leptin receptor are found in fish, the hormone is not expressed in adipose tissue, but is found in liver and other tissues. Here, we show that adult zebrafish lacking a functional leptin receptor do not exhibit hyperphagia or increased adiposity, and exhibit normal fertility. However, leptin receptor-deficient larvae have increased numbers of β -cells and increased levels of insulin mRNA. Furthermore, larval zebrafish have been shown to exhibit β -cell hyperplasia in response to high fat feeding or peripheral insulin resistance, and we show here that leptin receptor is required for this response. Adult zebrafish also have increased levels of insulin mRNA and other alterations in glucose homeostasis. Thus, a role for leptin in the regulation of β -cell mass and glucose homeostasis appears to be conserved across vertebrates, whereas its role as an adipostatic factor is likely to be a secondary role acquired during the evolution of mammals.

leptin | zebrafish | glucose homeostasis | adipostasis

The hormone leptin was identified in mammalian adipocytes (1) and well characterized in mice and humans as an adipostatic hormone. It is secreted into the serum in proportion to adipose mass and homeostatically regulates adipose mass primarily via binding to a distinct leptin receptor expressed in behavioral, endocrine, and autonomic control circuits in the central nervous system (2, 3). Failure of leptin signaling, due to mutations in leptin or leptin receptor genes, results in hyperphagia and hypometabolism to produce extreme obesity, diabetes, and infertility. Leptin and leptin receptor are highly conserved across mammalian species. Mouse and human leptin proteins are 83% identical, and leptin receptor proteins are 75% identical. However, the mammalian leptin and leptin receptor amino acid sequences are less well conserved with those of lower vertebrates. Indeed, the use of primary sequence homology failed to identify the leptin gene in fish or birds; chromosomal synteny was ultimately used to identify the gene in these vertebrate classes (4). For example, the zebrafish leptin protein is only 19% identical to the human protein.

Although the amino acid sequences are divergent, the basic structural features and intracellular signaling mechanisms of leptin and its receptor appear to be conserved throughout vertebrates (4). Furthermore, administration of mammalian leptin in birds and fish caused an anorexigenic effect, suggesting conservation of function of the leptin system (5). However, although leptin in mammals is predominantly expressed in adipose tissue (6), leptin expression in fish and birds appears to be negligible in adipose tissue (7). The expression profile varies widely, with many studies reporting the liver as a site of expression in fish (4). Interestingly, multiple studies report a rise in plasma leptin upon fasting (5), a response diametrically opposite to the leptin decrease observed in fasting

mice (8). In fish, hepatic leptin expression rises upon fasting (7) and the fish liver can secrete leptin (9). Deletion of the leptin receptor in medaka produced a modest increase in food intake, and juvenile but not adult growth, and a modest increase in visceral but not total body fat (10). These studies cast doubt as to the role of leptin as an adipostatic factor in fish and, thus, roles for leptin in nonmammalian vertebrates remains to be established. We report here on the role of leptin signaling in the zebrafish, *Danio rerio*, using a zebrafish mutant for the leptin receptor, and using clustered regularly interspaced short palindromic repeats (CRISPR) gene editing to mutate both leptin receptor and leptin genes.

Results

Mutation of the Leptin Receptor in Adult Zebrafish Has Limited Effect on Body Size, Weight, Adiposity, and Feeding. The zebrafish genome contains one leptin receptor gene (*lepr*) (11), and two leptin genes, *lepa* and *lepb* (12). To study the leptin system in teleosts, we obtained a zebrafish line expressing a mutant form of the leptin receptor (13). The *sa1508* allele, obtained through screening for mutations after *N*-ethyl-*N*-nitrosourea (ENU) mutagenesis, is a C > A nonsense mutation that leads to a premature stop codon after the start of the cytoplasmic domain of the receptor (Fig. S1A). This mutant is comparable to the nonsignaling truncated leptin receptor isoform in *db/db* mice (14) and rainbow trout (15).

Given the role of the mammalian leptin receptor in the adipostat, we first looked at size, weight, and body composition in the homozygous *lepr^{sa1508/sa1508}* mutant compared with wild-type (WT) sibling animals, all produced from *lepr^{sa1508/+}* heterozygous in-crosses, and raised as a mixed genotype population to prevent density effects. We measured standard length (SL) 5 d after fertilization (dpf), and SL and weight at 29 dpf, 41 dpf, and 100 dpf, and found no effect of genotype upon SL or weight (Fig. 1 A and B).

Significance

The hormone leptin homeostatically maintains long-term fat stores in mammals. Made by adipocytes in proportion to total adipose mass, leptin functions by regulating behavioral, autonomic, and endocrine circuits in the CNS to control energy intake and expenditure. As leptin signals nutritional sufficiency, it also acts as a gating factor for reproductive maturation and competence. Defective leptin signaling in mammals results in hyperphagia, obesity, diabetes, and infertility. Much less is known about leptin in nonmammalian vertebrates; however, the teleost leptin homologue is not primarily expressed in adipocytes. Here we show that zebrafish leptin is not required for adipostasis, food intake, or reproduction. However, we show here that, as in mammals, zebrafish leptin retains a role in the regulation of glucose homeostasis.

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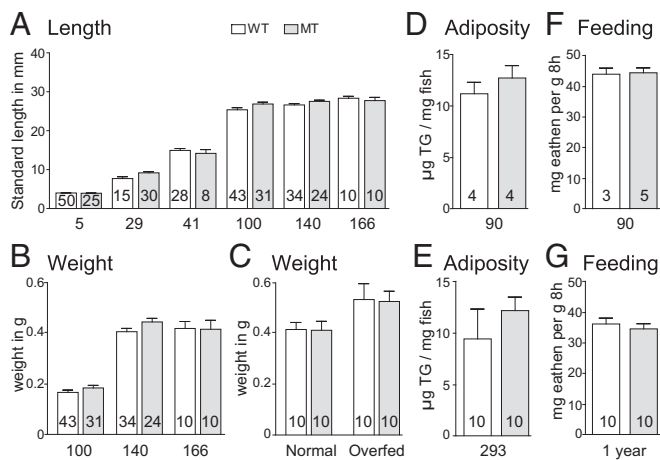


Fig. 1. Length, weight, adipose mass, and food intake in normal and leptin receptor mutant zebrafish. Heterozygous *lepr^{sa1508/+}* mutant fish were crossed to provide WT (open bars) and homozygous *lepr^{sa1508/1508}* (gray bars) offspring, which were grown together at similar densities, then characterized for growth, body composition, and feeding behavior, with the numbers (inside bars) and ages in days after fertilization (under bars) indicated. (A) Length of WT and *lepr^{sa1508/1508}* mutant fish from 5 to 166 dpf. Two-way ANOVA shows a significant effect of age [$F(5, 304) = 1,284, P < 0.0001$] but not genotype [$F(1, 304) = 2.104, P > 0.05$]. (B) Weight of WT and *lepr^{sa1508/1508}* mutant fish from 100 to 166 dpf. Two-way ANOVA shows a significant effect of age [$F(2, 151) = 227.1, P < 0.0001$] but not genotype [$F(1, 151) = 1.706, P > 0.05$]. (C) Weight of WT and *lepr^{sa1508/1508}* mutant fish following 30 d of high fat feeding. Two-way ANOVA shows significant increase in weight as a function of diet [$F(1, 39) = 4.687, P < 0.05$] but not of genotype [$F(1, 39) = 0.03255, P > 0.05$]. (D and E) Total triglyceride content of WT and *lepr^{sa1508/1508}* mutant fish at 90 and 293 d of age, respectively. There was no significant difference at either age $t(6) = 0.8631, P > 0.4213$ (D) and $t(18) = 0.9246, P > 0.3673$ (E); two-tailed t test. (F and G) Food consumption of 3-mo-old male (F) and 1 y old female WT and *lepr^{sa1508/1508}* mutant (G) fish over 16 trials. A repeated-measures ANOVA shows no significant effect of genotype in males [$F(1, 127) = 0.00476, P > 0.05$] or females [$F(1, 319) = 1.013, P > 0.05$]. Data shown as means \pm SEM.

We separated animals by genotype, keeping them at the same density and scored SL after maturity at 4 and 5.5 mo (140 and 166 dpf, respectively), with no differences observed. Stratifying the data by gender from 100 dpf onward also did not identify any effects of genotype. We also examined WT and homozygous *lepr^{sa1508/1508}* fish by using an overfeeding paradigm demonstrated to produce obesity in the zebrafish (16). As expected, the “overfed” fish gained a significant amount of weight compared with normal fed fish (Fig. 1C). However, we observed no significant effect of genotype. We next investigated whole body adiposity in 3-mo-old male fish and again saw no significant difference between genotypes (Fig. 1D). To ascertain that there was no long-term adiposity effect of the *lepr^{sa1508/1508}* genotype, we kept two clutches of fish until 293 dpf, but saw no significant effect of genotype (Fig. 1E). The leptin-deficient *ob/ob* and leptin receptor-deficient *db/db* mice also exhibit profound hyperphagia (1). We trained male animals to recognize a commercially available fish food (Betta Bits, TopFin) and tested food intake for 16 d with 8 h of exposure to excess food, each followed by a day of rest. There was no effect of genotype on food intake either in 3-mo or 1-y-old sibling animals (Fig. 1F and G).

Mutation of the Leptin Receptor in Zebrafish Has No Effect on Fertility. An additional phenotype of leptin deficiency in mice and humans is infertility (17, 18). We therefore investigated reproductive competency by crossing wild-type siblings as well as *lepr^{sa1508/1508}* mutant adults and scoring the productivity and efficiency of breeding events (Fig. 2). Reproductive productivity was determined by counting individual clutch sizes of five couples of each genotype with variable periods of separation (daily breeding, 3–4 d,

6 d or ≥ 8 d of separation). Clutch size increases based on the time of parental separation, but no effect of genotype on the number of fertilized eggs laid (Fig. 2A) or frequency of successful breedings (Fig. 2B) was observed.

Mutation of the Leptin Receptor in Larval Zebrafish Increased Insulin and Glucagon Gene Expression, and β -Cell Mass. The leptin receptor-deficient *diabetes (db/db)* mouse exhibits hyperglycemia by 3–4 wk of age (19). Hepatic mRNA levels for leptin have been shown to change upon fasting in zebrafish, common carp, Atlantic salmon, and Arctic charr (5), and recombinant leptin was shown to induce hepatic glucose mobilization in tilapia (20). Thus, we next sought to examine effects of leptin receptor deficiency on glucose homeostasis in the zebrafish (Fig. 3).

Total glucose levels were measured in zebrafish fry at 6 dpf, a timepoint at which fry have consumed their yolk. We found a small but significant increase in the glucose content of homozygous *lepr^{sa1508/1508}* mutant fry compared with WT fry (Fig. 3A). We next investigated insulin mRNA levels at 6 dpf by quantitative RT-PCR (qPCR). Zebrafish has two preproinsulin genes, *insa* and *insb*, due to a genome duplication event (21), with both isoforms being expressed in the pancreas (22). We found that *insa* but not *insb* is significantly up-regulated at 6 dpf in *lepr^{sa1508/1508}* mutant fry (Fig. 3B). We also looked at glucagon transcript levels and found that both *glucagon a* and *b* were up-regulated at 6 dpf in *lepr^{sa1508/1508}* fry (Fig. 3C).

db/db mice exhibit altered β -cell numbers and hyperglycemia (23, 24). We therefore crossed the *lepr^{sa1508}* allele into a zebrafish line carrying a β -cell marker [*Tg(-1.2ins:H2BmCherry)*] (25) and counted β -cells in 5-dpf fry (Fig. 3D). We found that the number of β -cells was significantly increased by $\sim 25\%$ in *lepr^{sa1508/1508}* animals compared with their WT and heterozygous siblings (Fig. 3E). To confirm that this phenotype is specific to defective leptin receptor function, we injected *Tg(-1.2ins:H2BmCherry)* embryos at the one-cell stage with Cas9 mRNA and guide RNA against the *lepr* gene to disrupt the *lepr* gene by using CRISPR (26). We used a mutagenic or a nonmutagenic control guide RNA against exons in *lepr*, and also controlled for DNA damage by using a guide RNA against the *tyrosinase* gene. Using a heteroduplex mobility shift assay, we confirmed mutagenesis in the *tyrosinase* gene and one of the two targeted sites in the *lepr* gene. We found a significant 14% increase in β -cells for the functional guide RNA against *lepr* but not in the groups injected with the *tyrosinase* guide RNA or the non-mutagenic *lepr* guide RNA, confirming results in *lepr^{sa1508/1508}* fish (Fig. 3F). Furthermore, we found that a guide RNA against *lepa*, but not *lepb*, also led to a 17% increase in the number of β -cells, further confirming a role for leptin signaling in regulation of β -cell mass in the larval zebrafish (Fig. 3F). We followed up on the

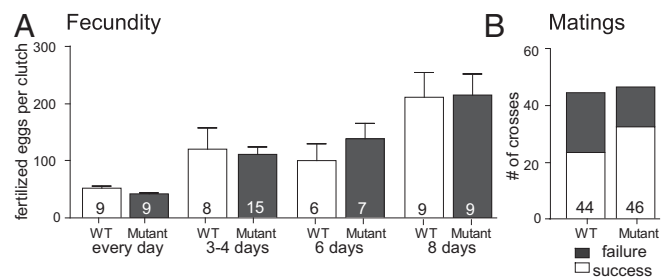


Fig. 2. Fecundity and mating efficiency in normal and leptin receptor mutant zebrafish. (A) Fertilized eggs from age-matched WT and *lepr^{sa1508/1508}* mutant couples bred daily or with variable times of separation between breeding, with days of separation indicated. Two-way ANOVA shows a significant effect for days of separation [$F(3, 71) = 13.29, P < 0.0001$] but not genotype [$F(1, 71) = 0.09, P > 0.05$] or an interaction of the two [$F(1, 71) = 0.29, P > 0.05$]. (B) Mating efficiency for WT and *lepr^{sa1508/1508}* mutant mating for fish of 3–6 mo of age. Data shown as means \pm SEM ($P > 0.05$, Fisher’s exact test); number of attempted matings indicated.

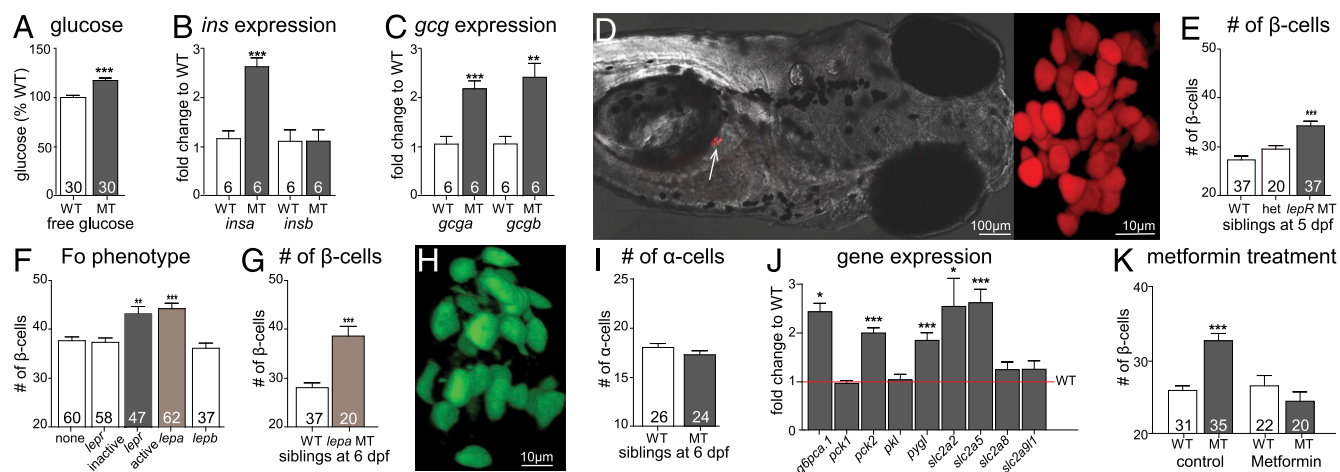


Fig. 3. Glucose homeostasis phenotypes in normal and leptin receptor mutant larval zebrafish. (A) Whole-body free glucose in WT and *lepr*^{sa1508/sa1508} mutant in-crosses collected at 6 dpf. A *t* test shows a significant difference between genotypes, $t(58) = 5.179$, $P < 0.0001$. (B) Expression of *insa* and *insb* mRNA at 6 dpf, determined by quantitative RT-PCR (qRT-PCR). *insa* shows significant up-regulation $t(10) = 6.058$, $P < 0.001$, whereas *insb* does not $t(10) = 0.016$, $P > 0.05$, *t* tests. (C) Expression of *gcga* and *gcgb* mRNA levels, determined by qRT-PCR. *gcga* $t(10) = 4.973$, $P < 0.001$ and *gcgb* $t(10) = 4.232$, $P < 0.01$ are up-regulated, *t* tests. (D) A representative image of a zebrafish carrying the β -cell marker [Tg(1.2ins:H2BmCherry)], and an enlarged image showing individual β -cells. (E) Numbers of β -cells in offspring of an in-cross of heterozygous *lepr*^{sa1508/sa1508} mutant animals at 5 dpf. Homozygous *lepr*^{sa1508/sa1508} mutants exhibit a significant increase in β -cell number by one-way ANOVA, $F(2, 93) = 18.69$, $P < 0.001$, Bonferroni's MCT. (F) Number of β -cells in animals at 6 dpf, following injection of embryos with Cas9 and guide RNA directed against *lepr*, *lepa*, or *lepb* genes; ANOVA $F(4, 263) = 11.67$, $P < 0.0001$, Bonferroni's MCT. (G) Number of β -cells is significantly increased in larvae homozygous for a mutation in *lepa* compared with their WT siblings at 6 dpf. *t* test $t(41) = 4.738$, $P < 0.001$. (H) A representative picture of α -cells in fish carrying the transgene [Tg(*gcg*:GFP)]. (I) Numbers of α -cells observed in WT and *lepr*^{sa1508/sa1508} mutant siblings from heterozygous *lepr*^{sa1508/sa1508} in-crosses at 6 dpf. There was no effect of genotype [*t* test $t(24) = 0.96$, $P = 0.34$]. (J) Data showing gene expression analysis of fry at 6 dpf compared with WT controls. *t* tests show an up-regulation of glucose 6 phosphatase (*g6pca.1*; $t(9) = 7.059$, $P < 0.001$), mitochondrial phosphoenolpyruvate carboxykinase (*pck2*; $t(9) = 8.601$, $P < 0.001$), liver glycogen phosphorylase (*pygl*; $t(9) = 4.423$, $P < 0.01$), glucose transporters 2 (*slc2a2*; $t(9) = 2.808$, $P < 0.05$) and 5 [*slc2a5*; $t(10) = 5.160$, $P < 0.001$], but not cytoplasmic phosphoenolpyruvate carboxykinase [*pck1*; $t(9) = 0.4822$, $P > 0.05$], liver pyruvate kinase (*pklr*; $t(10) = 0.2263$, $P > 0.05$), glucose transporters 8 [*slc2a8*; $t(10) = 0.903$, $P > 0.05$] or 9 [*slc2a9l1*; $t(10) = 1.31$, $P > 0.05$]. (K) Effect of metformin on the developmental increase in β -cell number. Two-factor ANOVA shows an effect of genotype [$F(1,107) = 4.785$, $P < 0.05$] and treatment [$F(1,107) = 12.56$, $P < 0.001$]. A Bonferroni post test showed a significant elevation in the DMSO-treated *lepr*^{sa1508/sa1508} mutant group. Data shown as means \pm SEM. * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$.

involvement of *lepa* by establishing germ-line founders carrying mutations in *lepa* and testing their offspring by using the β -cell marker [Tg(-1.2ins:H2BmCherry)]. We found an increase in the number of β -cells at 6 dpf by $\sim 38\%$ (Fig. 3G). We also crossed Tg(*gcga*:eGFP), a transgene driving GFP expression in α -cells (25), into *lepr*^{sa1508/sa1508} fish, to look for effects of *lepr* deficiency on α -cells (Fig. 3H). No significant effect of genotype on the number of α -cells at 6 dpf (Fig. 3I) was observed.

Mutation of the Leptin Receptor in Larval Zebrafish Alters Expression of Genes Involved in Hepatic Glucose Metabolism. Next, we investigated transcript levels of key enzymes in hepatic glucose metabolism at 6 dpf (Fig. 3J). Using qPCR, we found an up-regulation of mitochondrial but not cytosolic phosphoenolpyruvate carboxykinase (*pck2* and *pck1*, respectively) in *lepr*^{sa1508/sa1508} mutant fry. We tested for liver-specific transcripts of glycogen phosphorylase (*pygl*) and pyruvate kinase (*pklr*) and found the former to be up-regulated in *lepr*^{sa1508/sa1508} mutant fry. Lastly, we looked at glucose 6 phosphatase (*g6pca.1*), and glucose transporter (*glut*) isoforms expressed in the zebrafish liver (2, 5, 8, and 9a; ref. 27). We found a significant up-regulation of *g6pase*, *slc2a2*, and *slc2a5* in *lepr*^{sa1508/sa1508} mutant fry, but not *slc2a8* or *slc2a9l1* (Fig. 3J). We exposed larvae to metformin, a drug thought to have beneficial effects in diabetes patients because of effects on hepatic glucose homeostasis and insulin sensitivity (28). Exposure of larvae to metformin from 3 to 5 dpf completely abolished the increase seen in *lepr*^{sa1508/sa1508} mutant fry at 5 dpf (Fig. 3K).

Mutation of the Leptin Receptor in the Adult Zebrafish Leads to Altered Glucose Tolerance and Hepatic Gene Expression. We next investigated aspects of glucose homeostasis in adult zebrafish. We saw no differences in fasting blood glucose between WT and homozygous *lepr*^{sa1508/sa1508} fish. Challenging the fish with 0.5 mg

of glucose per gram of fish, however, we saw improved glucose disposal at 30 min in homozygous *lepr*^{sa1508/sa1508} fish (Fig. 4A), consistent with increased capacity for insulin release. The zebrafish β -cells are diffusely distributed along the intestine, whereas the primary islet in the pancreas is attached to the intestine. We thus examined *insa* and *gcga* transcript levels in whole visceral tissue, ~ 1 – 2 h after a meal of brine shrimp. We found a threefold higher expression of both genes in homozygous adult *lepr*^{sa1508/sa1508} fish relative to WT levels (Fig. 4B), paralleling larval results. We next tested hepatic transcript levels by using qPCR and found changes in *pygl*, *slc2a5*, and *slc2a9l1* but not *pck1*, *pck2*, *g6pca.1*, *pk1*, *slc2a2*, or *slc2a8* (Fig. 4C). Despite no evidence for elevated blood glucose or reduced glucose tolerance, we tested for a diabetic-like phenotype in adult fish, reduced wound healing. Streptozotocin-treated adult zebrafish have been shown to exhibit abnormalities in fin regeneration (29). We therefore clipped fins of 7-mo-old WT and homozygous *lepr*^{sa1508/sa1508} fish and imaged these fins after 3 d of growth. Regrowth of the lepidotrichial rays was impaired, indicative of a defect in wound healing (Fig. 4D).

Mutation of the Leptin Receptor in Larval Zebrafish Blocks Nutrient-Induced β -Cell Compensation. In zebrafish, nutrient excess leads to an increase in larval β -cell neogenesis (25). Because we saw a similar increase in the *lepr*^{sa1508/sa1508} mutant fry, we wished to test whether chronic nutrient excess would be ineffective in further increasing β -cell numbers in *lepr*^{sa1508/sa1508} mutant fry. First, we determined when, during development, leptin receptor deficiency led to an increase in β -cell mass. We saw a slight trend at 3 dpf and a significant increase in β -cell number at 4 dpf (Fig. 5A). By exposing the larvae to 5% egg yolk by volume (26.5% lipid and 15.8% protein content by weight) for 8 h at 5 dpf, a timepoint at which most of the larvae's own yolk is consumed, we

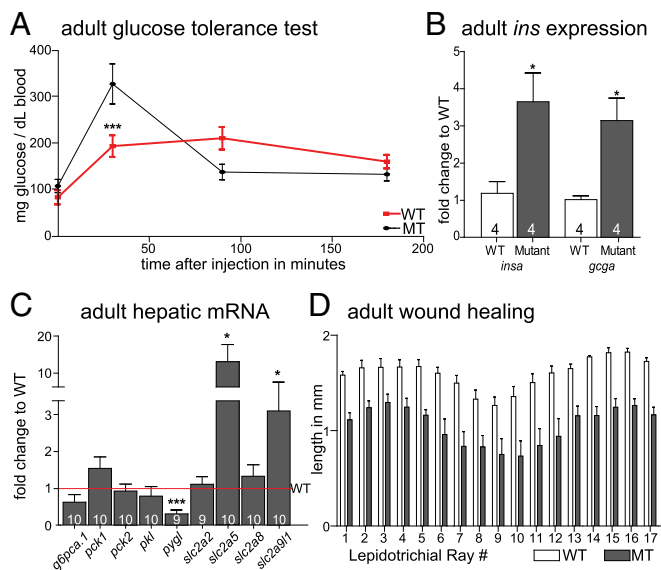


Fig. 4. Glucose homeostasis phenotypes in normal and leptin receptor mutant adult zebrafish. (A) Glucose tolerance test in 1-y-old WT and homozygous *lepr^{sa1508/sa1508}* fish. A two-way ANOVA post hoc test shows a significant decrease of blood glucose in homozygous *lepr^{sa1508/sa1508}* fish at 30 min with a significant interaction between time and genotype [$F(3,102) = 7.266$, $***P < 0.0001$]. (B) *insa* and *gcga* mRNA expression in adult visceral samples is increased [*insa* $t(6) = 2.947$, $*P < 0.05$ and *gcga* $t(6) = 3.459$, $*P < 0.05$]. (C) Adult hepatic gene expression analysis shows a down-regulation in liver glycogen phosphorylase [*pygl*; $t(17) = 4.650$, $***P < 0.001$], and up-regulation in glucose transporters 5 [*slc2a5*; $t(16) = 2.289$, $*P < 0.05$] and 9 [*slc2a9l1*; $t(17) = 2.616$, $*P < 0.05$] and no change in glucose 6 phosphatase [*g6pca.1*; $t(18) = 1.984$, $P = 0.0627$], cytoplasmic phosphoenolpyruvate carboxykinase [*pck1*; $t(17) = 0.3637$, $P = 0.72$], mitochondrial phosphoenolpyruvate carboxykinase [*pck2*; $t(18) = 0.6966$, $P = 0.49$], liver pyruvate kinase [*pklr*; $t(17) = 0.9630$, $P = 0.35$], glucose transporters 2 [*slc2a2*; $t(17) = 0.1563$, $P = 0.88$] or 8 [*slc2a8*; $t(18) = 0.9041$, $P = 0.38$], all t tests. (D) Tissue regeneration 3 d after fin-clipping. A two-way ANOVA shows a significant effect of genotype [$F(1, 186) = 169.7$, $P < 0.0001$]. Data shown as means \pm SEM.

found an increase in β -cell number in overfed wild-type animals but not in mutant *lepr^{sa1508/sa1508}* siblings. Indeed, nutrient excess seemed to equalize the number of β -cells between wild-type and homozygous *lepr^{sa1508/sa1508}* mutants (Fig. 5B). Persistent daily exposure to egg yolk leads to a further increase in β -cell mass during days 7–9 in WT larvae, but this late effect of nutrition was also completely blocked in *lepr^{sa1508/sa1508}* mutants (Fig. 5C). Thus, leptin receptor deficiency increases β -cell mass early in development, but at the same time, blocks nutrient-induced compensation. FGF1 mediates overnutrition-induced β -cell differentiation in the fish (30). We therefore tested whether an FGF signal mediates the early β -cell increase in *lepr^{sa1508/sa1508}* mutant larvae. We exposed larvae late at 3 dpf to SU5402 and scored β -cell number at 4 dpf. We found that the FGF inhibitor also blocks the early increase in β -cell number seen in *lepr^{sa1508/sa1508}* mutants (Fig. 5D), implicating FGF in the increase in β -cells resulting from either overnutrition or *lepr* deficiency.

Conclusions

The identification of leptin and leptin receptor gene homologs in lower vertebrates, such as birds and fish, has only recently allowed investigators to begin to address the comparative endocrinology of leptin action. A leptin receptor knockout reported in medaka (*Oryzias latipes*) (10) measured a 1.6- to twofold increase in post-juvenile body weight, but no increase in juvenile or adult body weight, and a 1.5- to 1.7-fold increase in postjuvenile and adult food intake. A small increase in visceral fat was reported, but without a detectable change in liver, muscle, or plasma triacylglycerol level. This increased adiposity is a relatively mild phenotype compared with leptin receptor signaling deficiency in mammals, which

exhibit severe hyperphagia, and morbid obesity in which total triacylglyceride levels can reach 50% of total body weight (31). For analysis of growth and body composition in *lepr* mutant zebrafish, we kept WT and mutant siblings of an in-cross of *lepr^{sa1508/+}* heterozygotes at the same density for half a year, and then scored weight and standard length at selected times. This approach, not used in the reported medaka experiments, is essential when working with fish to rule out effects of background genetics and tank densities. At no point did we see a difference in genotypes. Together with the limited phenotypes observed in medaka, these data argue that leptin does not act as an adipostatic factor in fish. In mammals, leptin has also acquired a role as a gating factor for reproductive competence, presumably informing the organism that adequate levels of energy are present to sustain a pregnancy. We found no difference in reproductive competency between WT and *lepr^{sa1508/sa1508}* mutant fish, arguing against a role for leptin in communicating information about long-term energy stores to reproductive circuits in the CNS.

Leptin has effects on glucose homeostasis in mice, independent of its effects on body weight or adiposity. For example, low-dose leptin can make *ob/ob* mice normoglycemic without affecting food intake or body weight (32). Remarkably, leptin has recently been demonstrated to normalize blood glucose, even in insulin-deficient mice (33). This regulatory action of leptin has been demonstrated to be mediated by leptin receptors in the CNS, and other effects have been reported in the periphery (for review, see ref. 34). Based on these findings, and our lack of evidence for adipostatic regulation by fish leptin, we examined effects of leptin signaling on glucose homeostasis in the zebrafish. When we looked at free glucose levels in fry, we observed a small increase in total body

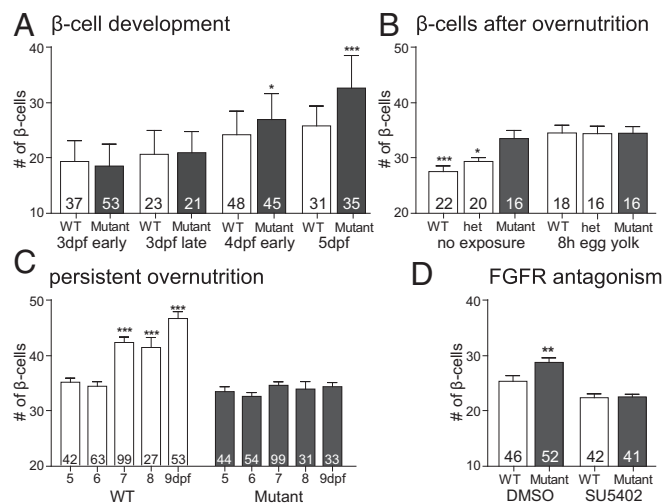


Fig. 5. β -cell development and regulation in normal and leptin receptor mutant zebrafish. (A) β -cell number in WT and *lepr^{sa1508/sa1508}* mutants develops between 3 and 5 dpf. A two-factor analysis showed a significant effect of genotype at 4 dpf [$F(1,254) = 6.4$, $P < 0.05$, Bonferroni post hoc test] and developmental time [$F(2, 254) = 59.83$, $P < 0.0001$]. (B) β -cell number increases following 8 h of high fat feeding (egg yolk) in 5 dpf animals produced from a *lepr^{sa1508/+}* heterozygous mutant in-cross [$F(1,106) = 19.38$, $P < 0.0001$, Bonferroni post hoc test]. *lepr^{sa1508/sa1508}* mutant animals already show an elevation in β -cell number and are nonresponsive to high-fat feeding [two-way ANOVA effect of genotype $F(2,106) = 3.3$, $P < 0.05$]. (C) Chronic high-fat feeding, with 8-h exposure to egg yolk each day from 5 to 9 dpf. Two-way ANOVA shows an effect of genotype [$F(1,544) = 79.72$, $P < 0.001$] and feeding [$F(4,544) = 14.59$, $P < 0.001$]. Bonferroni post hoc tests showed a compensatory β -cell increase in WT but not *lepr^{sa1508/sa1508}* animals. (D) Effect of FGFR inhibitor SU5402 on the developmental increase in β -cell number from 3 to 4 dpf. Two-factor ANOVA shows an effect of genotype [$F(1,180) = 4.935$, $P < 0.05$] as well as treatment [$F(1,180) = 32.73$, $P < 0.001$]. A Bonferroni post hoc test showed a significant elevation in the DMSO-treated *lepr^{sa1508/sa1508}* mutant group. Data shown as means \pm SEM. $*P < 0.05$; $**P < 0.01$; $***P < 0.001$.

glucose, along with elevated insulin and glucagon mRNA expression levels. It was, of course, not possible to measure blood glucose in larvae. We proceeded to score the number of α - and β -cells and found an early increase in the number of β -cells, but no difference in α -cells, in *lepr*^{sa1508/sa1508} mutant fry. In mice, lack of the leptin receptor leads to persistent hyperglycemia, hyperglucagonemia and hyperinsulinemia (35), concomitant with an increase in β -cell as well as α -cell mass (36). Both α - and β -cells are known to be able to significantly increase peptide hormone levels on a per-cell basis (37), thus there is clearly a differential response of these two islet cell types to a deficiency of leptin signaling in the zebrafish.

To confirm that these islet phenotypes resulted from defective leptin signaling, we used CRISPR mutagenesis in zebrafish embryos and characterized the resulting fry at a comparable age following mutagenesis of *lepr*, *lepa*, or *lepb* genes in a zebrafish line carrying a β -cell marker. Previous data show that this method can be used to characterize genotype–phenotype relationships in the F₀ generation, because biallelic mutagenesis at efficiencies of up to 80% can be readily achieved (26). We found that mutagenesis of *lepr* and *lepa*, but not *lepb*, exhibited increased β -cell number. We also replicated an increase in β -cell number in F₂ sibling larvae mutant for *lepa*, providing independent support of the validity of using CRISPR in the F₀ generation and the role of *lepr* and *lepa* in the islet phenotype. These data support the hypothesis that leptin signaling regulates β -cell mass in the larval zebrafish.

Genes involved in glucose metabolism in liver were also found to be dysregulated. We found that the mRNA for the mitochondrial (*pck2*) but not cytosolic form (*pck1*) of PEPCK is elevated. Although the cytosolic form has a more canonical role in gluconeogenesis, hepatic mPEPCK also plays a role in gluconeogenesis and silencing of the gene lowers blood glucose in mice (38). We furthermore saw an increase in hepatic glycogen phosphorylase expression (glycogenolysis) and no change in pyruvate kinase expression (glycolysis). Additionally, we observed an increase in glucose 6 phosphatase (*g6ca.1*) and glucose transporters 2 and 5. Taken together, these results are suggestive of increased gluconeogenesis and glycogenolysis in larvae. When we looked at adult hepatic expression ~5 h after a meal, however, we found transcript changes in glycogen phosphorylase and glucose transporters 5 and 9a, but none of the other transcripts. Together, the observed expression changes argue for a dysregulation in multiple glucose homeostasis pathways. Interestingly, metformin treatment normalized the number of β -cells in *lepr*^{sa1508/sa1508} mutant fry, providing some support for the idea that reduced leptin signaling exerts its effects through the liver and/or other peripheral tissues.

In contrast to larvae, adult fish exhibited fasting euglycemia, and improved glucose disposal after 30 min in a glucose tolerance test, suggestive of increased β -cell mass, but not insulin resistance. *insa* and *gcga* expression were increased in *lepr*^{sa1508/sa1508} mutant adults, paralleling the finding in larvae. These findings are similar to mouse models in which growth factor expression is targeted to β -cells. In these models, increased β -cell number and improved glucose tolerance are seen in the absence of obesity or insulin resistance (e.g., ref. 39). Mice with either liver- or pancreas-specific knockout of *lepr* show a similar phenotype, with euglycemia, and improved glucose clearance in a GTT due to an enhanced early phase insulin secretion (40, 41). The liver-specific *lepr* knockout mice also show an increase in β -cell mass (41). A speculative suggestion is that the conserved function of leptin seen in the fish parallels that documented for peripheral leptin action in the mouse: regulation of islet development and function. Despite the absence of obesity or glucose intolerance in adult *lepr*^{sa1508/sa1508} fish, these animals exhibited a defect in wound healing, a finding previously validated in a fish model of diabetes (29). Additional work will be needed to determine whether this is a true diabetic phenotype or altered function of one or more growth factors unrelated to blood glucose levels.

High-fat feeding in zebrafish leads to a compensatory increase in the number of β -cells (25). A similar response to defective

Table 1. Comparison of leptin action in mammals and larval zebrafish

Phenotype	Mammal	Zebrafish
Hyperphagia	Yes	No
Hypometabolism	Yes	Not tested
Obesity	Yes	No
Hyperinsulinemia	Yes	Yes*
Hyperglucagonemia	Yes	Yes*
Diabetes	Yes	Yes [†]
Infertility	Yes	No
Immune dysfunction	Yes	Not tested

*Suggested by increased total gene expression.

[†]In larvae.

leptin signaling, shown here, suggested that the nutritional signal leading to an increase in β -cell number may require leptin. Indeed, *lepr*^{sa1508/sa1508} animals did not respond to either an acute or sustained nutrient challenge by increasing the number of β -cells. This result suggests that *lepr*^{sa1508/sa1508} mutant animals lack a signaling pathway critical for compensatory increase in β -cell mass in response to overnutrition. Data in the zebrafish (30) show that nutrient excess leads β -cells to secrete FGF, which then leads to β -cell differentiation. The FGF receptor (FGFR) antagonist SU5402 also blocks the increase in β -cell number due to defective leptin signaling. Therefore, FGF signaling appears to act downstream of leptin in the regulation of β -cell mass.

In conclusion, the data reported here (Table 1), particularly in the context of the limited expression of leptin in adipose tissue in fish, support the hypothesis that leptin is not an adipostatic factor in fish. The data show a sustained effect of *lepr* deficiency on total insulin and glucagon transcript levels, and dysregulated hepatic gene expression in larval and adult fish. Data from larvae shows clear effects on developmental and nutritional regulation of β -cell number. Further studies are needed to determine effects of zebrafish leptin on hepatic glucose production, glucose action, and insulin action. Because regulation of components of glucose homeostasis appears to be a conserved function of leptin in fish and mammals, these data suggest that this function has been conserved throughout vertebrate evolution, and that the role of leptin in adipostasis developed subsequently in mammals, or was lost and supplanted by an as-yet unknown signaling pathway in fish.

Methods

Zebrafish Strains and Maintenance. Larvae were raised in 28 °C incubators on 14:10h light-dark cycles until 5 dpf when they were placed on a standard diet and maintained in an Aquatic Habitats system on a 14:10h light-dark cycle. The *lepr* mutant strain *sa1508* was obtained from the Zebrafish International Resource Center. *Tg(-1.2ins:H2BmCherry)* and *Tg(gcga:eGFP)* were obtained from the W.C. laboratory (25). The *lepa* mutant line was established by using a gRNA to the target (AATCTCTGGATAATGTCTCTGG) as described (26). All procedures were approved by the Vanderbilt Institutional Animal Care and Use Committee.

Body Composition. Animals were isolated, anesthetized with tricaine solution and standard length measured. Animals were then blotted dry, weighed, and allowed to recover before being returned to their home tank. Whole-body adiposity was analyzed by the Vanderbilt Hormone Assay and Analytical Services core. For all body composition experiments, animals were kept at the same tank density (42).

Feeding. Standard feeding protocol was two meals of live artemia and two meals of Tetra-Min flakes (Tetra). For overfeeding, control fish were fed a reduced diet of one meal of artemia a day, whereas overfed animals were fed ad libitum artemia from approximately 9:00 AM until about 4:00 PM. Uneaten and/or dead artemia were removed before each feeding. For food consumption, animals were fed betta bit pellets (TopFin), exclusively off the system with frequent water changes for a month. Fish were then weighed and individually kept in 3-L tanks. Animals were fed every other day with

excess food pellets, and the remaining pellets were counted and removed after 8 h of overfeeding with yolk, as described (25).

RNA Extraction and qPCR. RNA was extracted from pools of 10 fry for larvae and from dissected male viscera or liver for adults by using TRIzol (Invitrogen). RNA was measured, and 1 μg of RNA with a 260/280 ratio >1.8 was used for reverse transcription by using a high-capacity cDNA kit (ABI). Gene expression was measured with SYBR green (ABI/Promega) by using a CFX96 cyclor (Bio-Rad). Relative expression compared with WT was calculated based on primer efficiency by using a pooled sample of the relative dataset's cDNA, then standardized to *eef1a11* expression. Primer sequences are sent upon request.

α - and β -Cell Counting. As described (25), representative pictures were taken with a Zeiss LSM710 META by using the Vanderbilt Cell Imaging Shared Resource core.

F₀ CRISPR Experiment. Offspring of *Tg(-1.2ins:H2BmCherry)* animals were injected at the one-cell stage into the cell with ~ 1 nL of a solution containing zebrafish 150 $\mu\text{g}/\mu\text{L}$ Cas9 mRNA (26) and 125 $\mu\text{g}/\mu\text{L}$ gRNA. Efficiency was analyzed by using heteroduplex mobility assay analysis. Target sequences were *lepr* (inactive) AGCATGATGAAGACAGACCTAGG; *lepr* (active) GGAGCGCCAGTAAAGCCGTGTGG; *lepa* GGAATCTCTGGATAATGTCCTGG, *lepb* ACAGAAGTACGACCATCAATGGG; and *tyr* (26).

Free Glucose Assay. Whole-body glucose was determined by using an Amplex Red glucose assay kit (Life Technologies). Briefly, sets of five fry were homogenized

as a pool in sample buffer on ice, the homogenate was cleared, and the supernatant measured according to the included instructions.

Glucose Tolerance Test. The test was performed according to refs. 43 and 44. Briefly, zebrafish were starved overnight, anesthetized with 3-aminobenzoic acid ethyl ester methanesulfonate, and injected i.p. with 0.5 mg of glucose per gram of fish weight in Hanks' buffered salt solution (Cellgro). Blood was collected from the tail and immediately diluted with assay buffer and snap frozen until measured (Amplex Red, Life Technologies).

Fin Regeneration. Fin regeneration was carried out as described in ref. 29. Briefly, fins were cut proximal to the proximal branch point of the dermal rays and allowed to recover for 3 d in the system before a picture was taken. Growth from the cut site was measured using ImageJ.

Compound Exposure. SU5402 (Tocris) was dissolved in DMSO and added to 0.3 \times Danieau solution at a final concentration of 15 μM for 24 h from 3 dpf to 4 dpf (30). Metformin (Sigma) was dissolved to 100 mM in 0.3 \times Danieau solution and used at 250 μM (45), with exposure from 3 dpf until 5 dpf. Solution was changed twice daily.

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